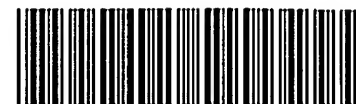


RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/630,518 A
Source: FW16
Date Processed by STIC: 02/08/2006

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 02/08/2006

PATENT APPLICATION: US/10/630,518A

TIME: 12:08:12

Input Set : A:\SEQ LIST 532792001100.txt

Output Set: N:\CRF4\02082006\J630518A.raw

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4 <110> APPLICANT: THE SALK INSTITUTE FOR BIOLOGICAL STUDIES
5     THE REGENTS OF THE UNIVERSITY OF CALIFORNIA
6     LILJEGREN, Sarah, Jean
7     ECKER, Joseph, R.
8     YANOFSKY, Martin, F.
10 <120> TITLE OF INVENTION: GENETIC CONTROL OF ORGAN ABSCISSION
13 <130> FILE REFERENCE: 532792001100
15 <140> CURRENT APPLICATION NUMBER: 10/630,518A
16 <141> CURRENT FILING DATE: 2003-07-29
18 <150> PRIOR APPLICATION NUMBER: US 60/264,974
19 <151> PRIOR FILING DATE: 2001-01-29
21 <150> PRIOR APPLICATION NUMBER: PCTUS02/01938
22 <151> PRIOR FILING DATE: 2002-01-22
24 <160> NUMBER OF SEQ ID NOS: 26
26 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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30 <212> TYPE: DNA
31 <213> ORGANISM: Arabidopsis thaliana
33 <220> FEATURE:
34 <221> NAME/KEY: CDS
35 <222> LOCATION: (1)...(1452)
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40 1 5 10 15
42 aag att ctt gaa ggg ctt ctt aaa cat cca gag aac aga gaa tgt gct 96
43 Lys Ile Leu Glu Gly Leu Leu Lys His Pro Glu Asn Arg Glu Cys Ala
44 20 25 30
46 gac tgc aaa aca aaa ggt cca aga tgg gct agt gtt aat tta ggt atc 144
47 Asp Cys Lys Thr Lys Gly Pro Arg Trp Ala Ser Val Asn Leu Gly Ile
48 35 40 45
50 ttt atc tgc atg caa tgt tct ggg att cac agg agt ctc ggg gta cac 192
51 Phe Ile Cys Met Gln Cys Ser Gly Ile His Arg Ser Leu Gly Val His
52 50 55 60
54 ata tcg aag gtt cga tct gcc act ctg gac aca tgg ctc ccc gag cag 240
55 Ile Ser Lys Val Arg Ser Ala Thr Leu Asp Thr Trp Leu Pro Glu Gln
56 65 70 75 80
58 gtt gca ttt ata cag tca atg gga aat gat aaa gca aat agt tac tgg 288
59 Val Ala Phe Ile Gln Ser Met Gly Asn Asp Lys Ala Asn Ser Tyr Trp
60 85 90 95
62 gaa gca gag cta ccc cca aac tat gat aga gtt gga att gag aat ttt 336
63 Glu Ala Glu Leu Pro Pro Asn Tyr Asp Arg Val Gly Ile Glu Asn Phe

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TIME: 12:08:12

Input Set : A:\SEQ LIST 532792001100.txt

Output Set: N:\CRF4\02082006\J630518A.raw

64	100	105	110	
66	ata cgt gca aag tat gaa gag aag aga tgg gtt tct aga ggg gaa aag	384		
67	Ile Arg Ala Lys Tyr Glu Glu Lys Arg Trp Val Ser Arg Gly Glu Lys			
68	115	120	125	
70	gct aga tca cct cct aga gtc gag cag gaa cgg cgg aaa tct gtg gag	432		
71	Ala Arg Ser Pro Pro Arg Val Glu Gln Glu Arg Arg Lys Ser Val Glu			
72	130	135	140	
74	aga agt ggg ccg gga tat gag cat gga cat agt agt agt cct gta aat	480		
75	Arg Ser Gly Pro Gly Tyr Glu His Gly His Ser Ser Ser Pro Val Asn			
76	145	150	155	160
78	ttg ttt gag gag agg aaa act att cca gca tct aga aca aga aat aat	528		
79	Leu Phe Glu Glu Arg Lys Thr Ile Pro Ala Ser Arg Thr Arg Asn Asn			
80	165	170	175	
82	gtt gct gca acg aga ata aat ctt ccc gtg cct ccc caa gga ccc agt	576		
83	Val Ala Ala Thr Arg Ile Asn Leu Pro Val Pro Pro Gln Gly Pro Ser			
84	180	185	190	
86	cag gtt ata aag cca cag cag aaa atg gag tct gca gct act cca gta	624		
87	Gln Val Ile Lys Pro Gln Gln Lys Met Glu Ser Ala Ala Thr Pro Val			
88	195	200	205	
90	gag agg gag aaa caa gca gta aat gtt gca cca gca tca gat cct cca	672		
91	Glu Arg Glu Lys Gln Ala Val Asn Val Ala Pro Ala Ser Asp Pro Pro			
92	210	215	220	
94	aag gtg gat ttt gct act gat ctg ttt aac atg cta tca atg gat gat	720		
95	Lys Val Asp Phe Ala Thr Asp Leu Phe Asn Met Leu Ser Met Asp Asp			
96	225	230	235	240
98	tcg act aca aat acc tca gag gca act cct ggc gat act cct gcc gat	768		
99	Ser Thr Thr Asn Thr Ser Glu Ala Thr Pro Gly Asp Thr Pro Ala Asp			
100	245	250	255	
102	gat aac tca tgg gct ggc ttt cag tct gct gga agt ggt caa acg gca	816		
103	Asp Asn Ser Trp Ala Gly Phe Gln Ser Ala Gly Ser Gly Gln Thr Ala			
104	260	265	270	
106	gag aaa att gtc aca gcc aag cct gct gag agc agt tct cct cca gct	864		
107	Glu Lys Ile Val Thr Ala Lys Pro Ala Glu Ser Ser Ser Pro Pro Ala			
108	275	280	285	
110	tca tct tct gac ttt gag gat ttg ttt aag gac aca cct aat tta aca	912		
111	Ser Ser Ser Asp Phe Glu Asp Leu Phe Lys Asp Thr Pro Asn Leu Thr			
112	290	295	300	
114	act caa caa gca cca aaa gat gtg aaa ggc gat atc atg agc ctg ttt	960		
115	Thr Gln Gln Ala Pro Lys Asp Val Lys Gly Asp Ile Met Ser Leu Phe			
116	305	310	315	320
118	gag aag acg aat ata gta tcg cct ttt gcc atg cat cag caa cag gtt	1008		
119	Glu Lys Thr Asn Ile Val Ser Pro Phe Ala Met His Gln Gln Gln Val			
120	325	330	335	
122	gct atg ctc gct cag cag caa gcc ctt tac atg gct gca gcg aaa gct	1056		
123	Ala Met Leu Ala Gln Gln Gln Ala Leu Tyr Met Ala Ala Ala Lys Ala			
124	340	345	350	
126	gct gga ggc act cca aac ggc gtg aat caa caa gct att gct aat gct	1104		
127	Ala Gly Gly Thr Pro Asn Gly Val Asn Gln Gln Ala Ile Ala Asn Ala			
128	355	360	365	

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Input Set : A:\SEQ LIST 532792001100.txt

Output Set: N:\CRF4\02082006\J630518A.raw

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130 ctt aac gta gct tct gca aat tgg tca aac ccc ggc ggc tac cag atc 1152
131 Leu Asn Val Ala Ser Ala Asn Trp Ser Asn Pro Gly Gly Tyr Gln Ile
132 370 375 380
134 ccc gga atg act aac ccc gta ggt ggt caa gct gat ctc cag aaa ctt 1200
135 Pro Gly Met Thr Asn Pro Val Gly Gly Gln Ala Asp Leu Gln Lys Leu
136 385 390 395 400
138 atg caa aac atg aat atg aac gca aac atg aac acg aga ccc gca caa 1248
139 Met Gln Asn Met Asn Met Asn Ala Asn Met Asn Thr Arg Pro Ala Gln
140 405 410 415
142 ccg caa gag aac act cta caa tac cca tca tcc agt ttc tac aca atg 1296
143 Pro Gln Glu Asn Thr Leu Gln Tyr Pro Ser Ser Ser Phe Tyr Thr Met
144 420 425 430
146 ggt caa gct aat caa gtg aac ggt atg acc cca aac tca acc ggt aaa 1344
147 Gly Gln Ala Asn Gln Val Asn Gly Met Thr Pro Asn Ser Thr Gly Lys
148 435 440 445
150 cct cag tca tca tcc gca acc caa cca aca agc acc aca cca tct tca 1392
151 Pro Gln Ser Ser Ser Ala Thr Gln Pro Thr Ser Thr Thr Pro Ser Ser
152 450 455 460
154 caa tca ggc aaa gac ttt gat ttc tct tcc ttg atg gat gga atg ttc 1440
155 Gln Ser Gly Lys Asp Phe Asp Phe Ser Ser Leu Met Asp Gly Met Phe
156 465 470 475 480
158 aca aaa cat tga 1452
159 Thr Lys His *
163 <210> SEQ ID NO: 2
164 <211> LENGTH: 483
165 <212> TYPE: PRT
166 <213> ORGANISM: Arabidopsis thaliana
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171 Lys Ile Leu Glu Gly Leu Leu Lys His Pro Glu Asn Arg Glu Cys Ala
172 20 25 30
173 Asp Cys Lys Thr Lys Gly Pro Arg Trp Ala Ser Val Asn Leu Gly Ile
174 35 40 45
175 Phe Ile Cys Met Gln Cys Ser Gly Ile His Arg Ser Leu Gly Val His
176 50 55 60
177 Ile Ser Lys Val Arg Ser Ala Thr Leu Asp Thr Trp Leu Pro Glu Gln
178 65 70 75 80
179 Val Ala Phe Ile Gln Ser Met Gly Asn Asp Lys Ala Asn Ser Tyr Trp
180 85 90 95
181 Glu Ala Glu Leu Pro Pro Asn Tyr Asp Arg Val Gly Ile Glu Asn Phe
182 100 105 110
183 Ile Arg Ala Lys Tyr Glu Glu Lys Arg Trp Val Ser Arg Gly Glu Lys
184 115 120 125
185 Ala Arg Ser Pro Pro Arg Val Glu Gln Glu Arg Arg Lys Ser Val Glu
186 130 135 140
187 Arg Ser Gly Pro Gly Tyr Glu His Gly His Ser Ser Ser Pro Val Asn
188 145 150 155 160
189 Leu Phe Glu Glu Arg Lys Thr Ile Pro Ala Ser Arg Thr Arg Asn Asn

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TIME: 12:08:12

Input Set : A:\SEQ LIST 532792001100.txt

Output Set: N:\CRF4\02082006\J630518A.raw

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190          165          170          175
191 Val Ala Ala Thr Arg Ile Asn Leu Pro Val Pro Pro Gln Gly Pro Ser
192          180          185          190
193 Gln Val Ile Lys Pro Gln Gln Lys Met Glu Ser Ala Ala Thr Pro Val
194          195          200          205
195 Glu Arg Glu Lys Gln Ala Val Asn Val Ala Pro Ala Ser Asp Pro Pro
196          210          215          220
197 Lys Val Asp Phe Ala Thr Asp Leu Phe Asn Met Leu Ser Met Asp Asp
198 225          230          235          240
199 Ser Thr Thr Asn Thr Ser Glu Ala Thr Pro Gly Asp Thr Pro Ala Asp
200          245          250          255
201 Asp Asn Ser Trp Ala Gly Phe Gln Ser Ala Gly Ser Gly Gln Thr Ala
202          260          265          270
203 Glu Lys Ile Val Thr Ala Lys Pro Ala Glu Ser Ser Ser Pro Pro Ala
204          275          280          285
205 Ser Ser Ser Asp Phe Glu Asp Leu Phe Lys Asp Thr Pro Asn Leu Thr
206          290          295          300
207 Thr Gln Gln Ala Pro Lys Asp Val Lys Gly Asp Ile Met Ser Leu Phe
208 305          310          315          320
209 Glu Lys Thr Asn Ile Val Ser Pro Phe Ala Met His Gln Gln Gln Val
210          325          330          335
211 Ala Met Leu Ala Gln Gln Gln Ala Leu Tyr Met Ala Ala Ala Lys Ala
212          340          345          350
213 Ala Gly Gly Thr Pro Asn Gly Val Asn Gln Gln Ala Ile Ala Asn Ala
214          355          360          365
215 Leu Asn Val Ala Ser Ala Asn Trp Ser Asn Pro Gly Gly Tyr Gln Ile
216          370          375          380
217 Pro Gly Met Thr Asn Pro Val Gly Gly Gln Ala Asp Leu Gln Lys Leu
218 385          390          395          400
219 Met Gln Asn Met Asn Met Asn Ala Asn Met Asn Thr Arg Pro Ala Gln
220          405          410          415
221 Pro Gln Glu Asn Thr Leu Gln Tyr Pro Ser Ser Ser Phe Tyr Thr Met
222          420          425          430
223 Gly Gln Ala Asn Gln Val Asn Gly Met Thr Pro Asn Ser Thr Gly Lys
224          435          440          445
225 Pro Gln Ser Ser Ser Ala Thr Gln Pro Thr Ser Thr Thr Pro Ser Ser
226          450          455          460
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229 Thr Lys His
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234 <211> LENGTH: 1452
235 <212> TYPE: DNA
236 <213> ORGANISM: Arabidopsis thaliana
238 <220> FEATURE:
239 <221> NAME/KEY: CDS
240 <222> LOCATION: (1)...(1452)
242 <400> SEQUENCE: 3
243 atg aac gag aaa gcc aac gtc tct aag gag ctt aat gcc cgc cat aga 48

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RAW SEQUENCE LISTING

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Input Set : A:\SEQ LIST 532792001100.txt

Output Set: N:\CRF4\02082006\J630518A.raw

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245 1 5 10 15
247 aag att ctt gaa ggg ctt ctt aaa cat cca gag aac aga gaa tgt gct 96
248 Lys Ile Leu Glu Gly Leu Leu Lys His Pro Glu Asn Arg Glu Cys Ala
249 20 25 30
251 gac tgc aaa aca aaa ggt cca aga tgg gct agt gtt aat tta ggt atc 144
252 Asp Cys Lys Thr Lys Gly Pro Arg Trp Ala Ser Val Asn Leu Gly Ile
253 35 40 45
255 ttt atc tac atg caa tgt tct ggg att cac agg agt ctc ggg gta cac 192
256 Phe Ile Tyr Met Gln Cys Ser Gly Ile His Arg Ser Leu Gly Val His
257 50 55 60
259 ata tcg aag gtt cga tct gcc act ctg gac aca tgg ctc ccc gag cag 240
260 Ile Ser Lys Val Arg Ser Ala Thr Leu Asp Thr Trp Leu Pro Glu Gln
261 65 70 75 80
263 gtt gca ttt ata cag tca atg gga aat gat aaa gca aat agt tac tgg 288
264 Val Ala Phe Ile Gln Ser Met Gly Asn Asp Lys Ala Asn Ser Tyr Trp
265 85 90 95
267 gaa gca gag cta ccc cca aac tat gat aga gtt gga att gag aat ttt 336
268 Glu Ala Glu Leu Pro Pro Asn Tyr Asp Arg Val Gly Ile Glu Asn Phe
269 100 105 110
271 ata cgt gca aag tat gaa gag aag aga tgg gtt tct aga ggg gaa aag 384
272 Ile Arg Ala Lys Tyr Glu Glu Lys Arg Trp Val Ser Arg Gly Glu Lys
273 115 120 125
275 gct aga tca cct cct aga gtc gag cag gaa cgg cgg aaa tct gtg gag 432
276 Ala Arg Ser Pro Pro Arg Val Glu Gln Glu Arg Arg Lys Ser Val Glu
277 130 135 140
279 aga agt ggg ccg gga tat gag cat gga cat agt agt agt cct gta aat 480
280 Arg Ser Gly Pro Gly Tyr Glu His Gly His Ser Ser Ser Pro Val Asn
281 145 150 155 160
283 ttg ttt gag gag agg aaa act att cca gca tct aga aca aga aat aat 528
284 Leu Phe Glu Glu Arg Lys Thr Ile Pro Ala Ser Arg Thr Arg Asn Asn
285 165 170 175
287 gtt gct gca acg aga ata aat ctt ccc gtg cct ccc caa gga ccc agt 576
288 Val Ala Ala Thr Arg Ile Asn Leu Pro Val Pro Pro Gln Gly Pro Ser
289 180 185 190
291 cag gtt ata aag cca cag cag aaa atg gag tct gca gct act cca gta 624
292 Gln Val Ile Lys Pro Gln Gln Lys Met Glu Ser Ala Ala Thr Pro Val
293 195 200 205
295 gag agg gag aaa caa gca gta aat gtt gca cca gca tca gat cct cca 672
296 Glu Arg Glu Lys Gln Ala Val Asn Val Ala Pro Ala Ser Asp Pro Pro
297 210 215 220
299 aag gtg gat ttt gct act gat ctg ttt aac atg cta tca atg gat gat 720
300 Lys Val Asp Phe Ala Thr Asp Leu Phe Asn Met Leu Ser Met Asp Asp
301 225 230 235 240
303 tcg act aca aat acc tca gag gca act cct ggc gat act cct gcc gat 768
304 Ser Thr Thr Asn Thr Ser Glu Ala Thr Pro Gly Asp Thr Pro Ala Asp
305 245 250 255
307 gat aac tca tgg gct ggc ttt cag tct gct gga agt ggt caa acg gca 816
308 Asp Asn Ser Trp Ala Gly Phe Gln Ser Ala Gly Ser Gly Gln Thr Ala

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VERIFICATION SUMMARY

DATE: 02/08/2006

PATENT APPLICATION: US/10/630,518A

TIME: 12:08:13

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